

FIG. 1

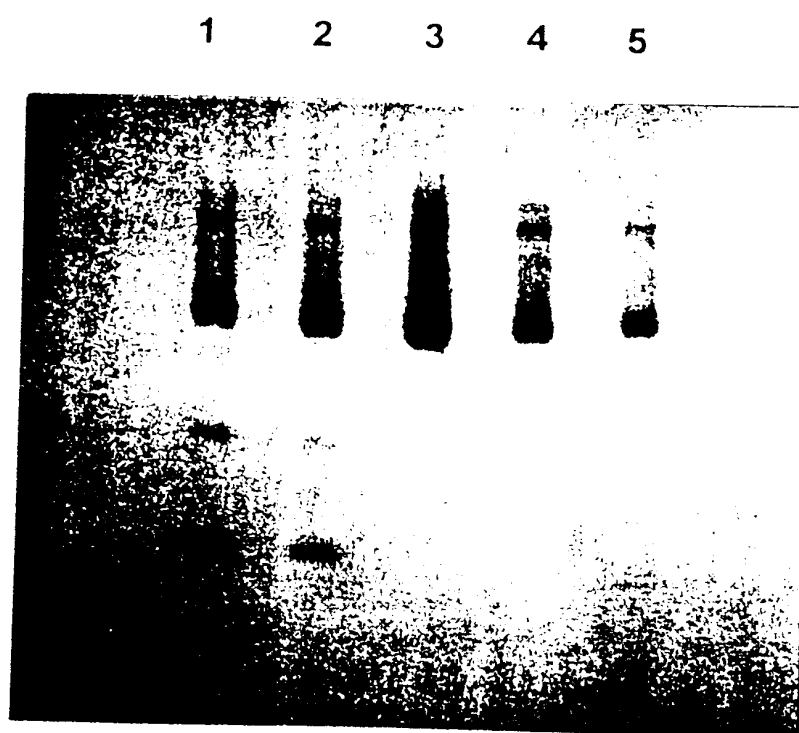


FIG. 2

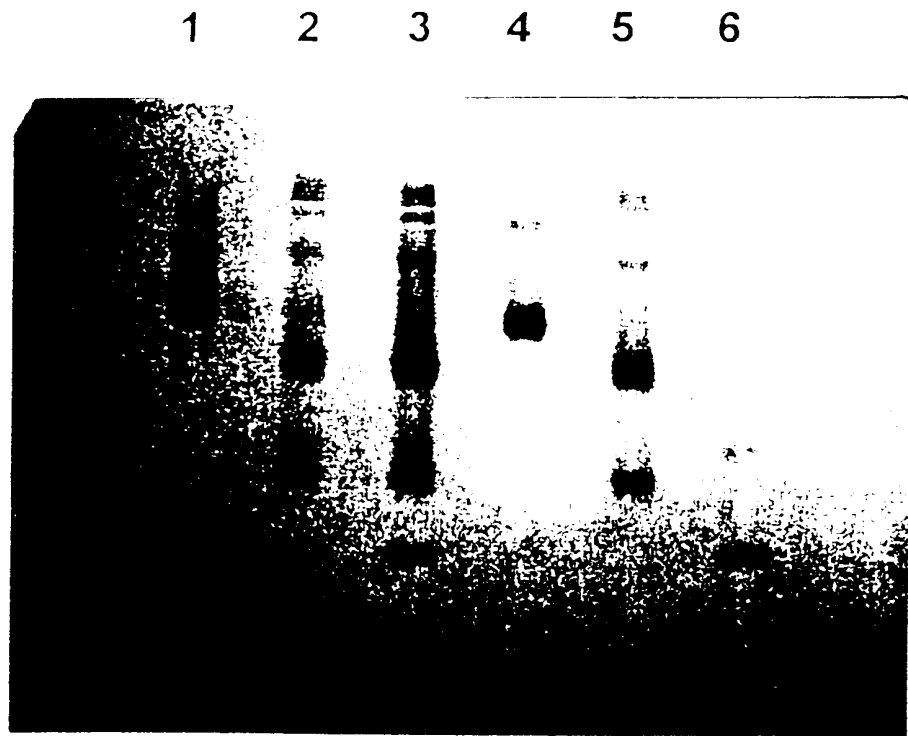


FIG. 3

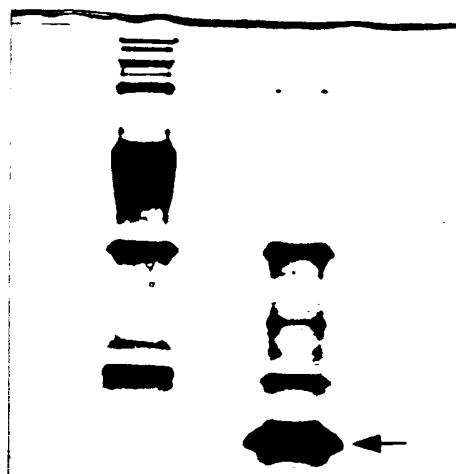


FIG. 4

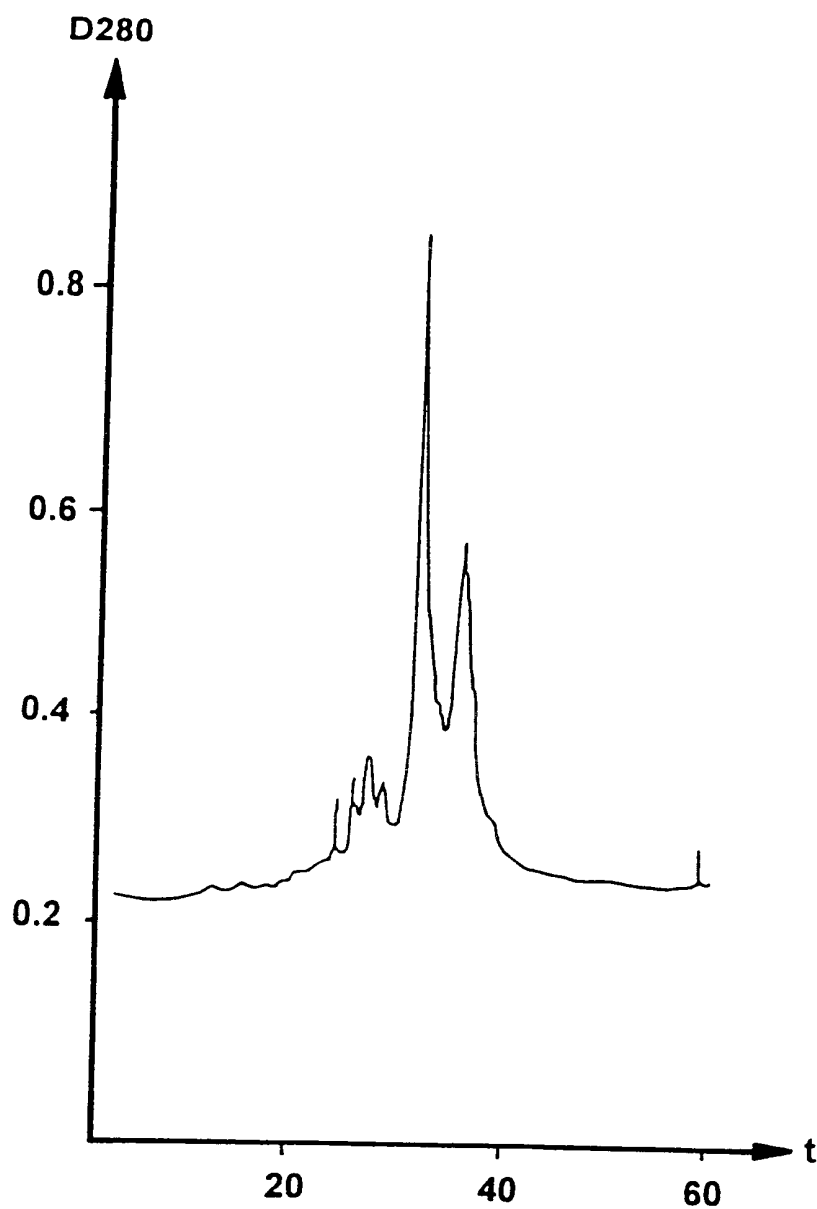


FIG. 5

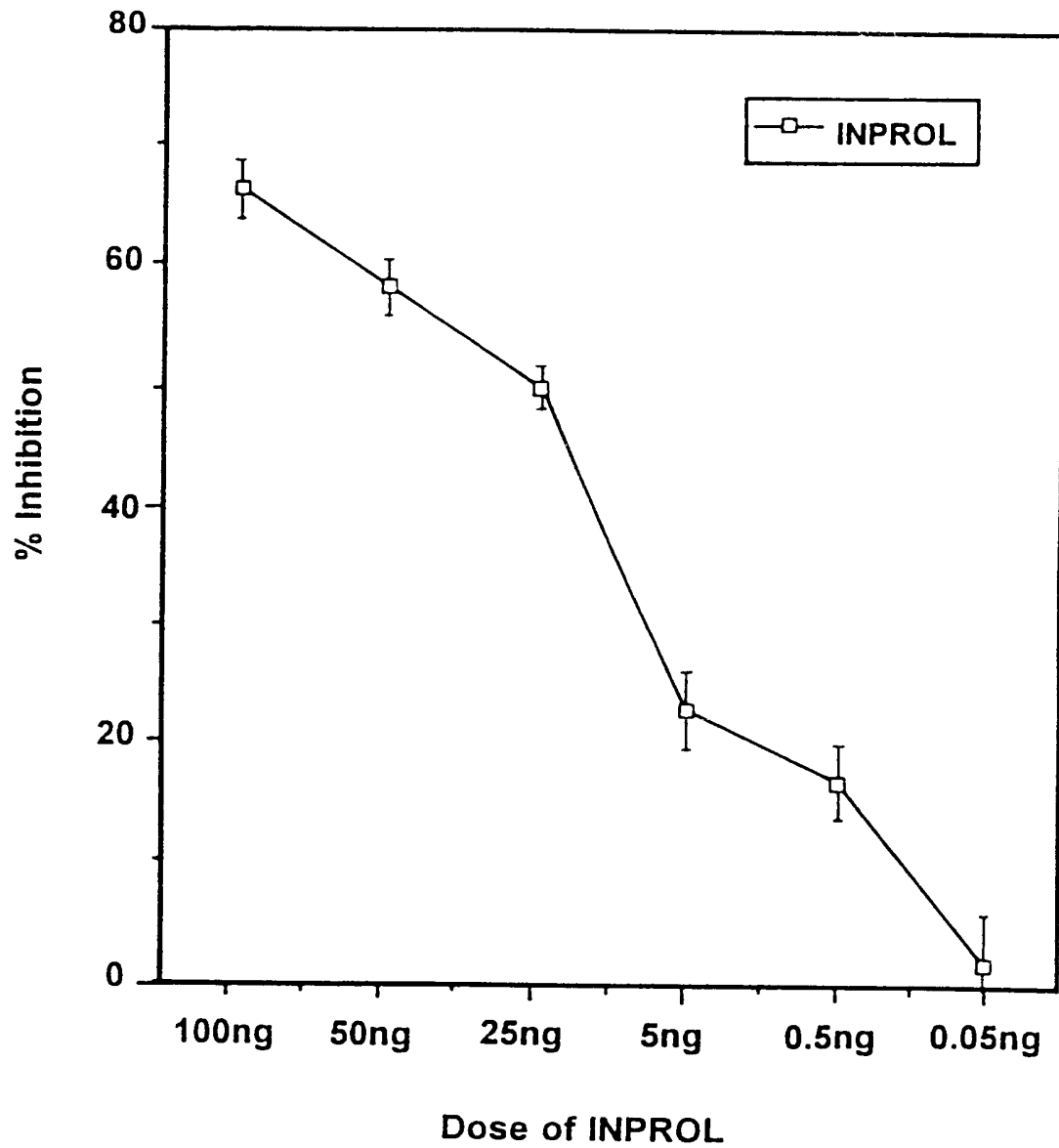


FIG. 6

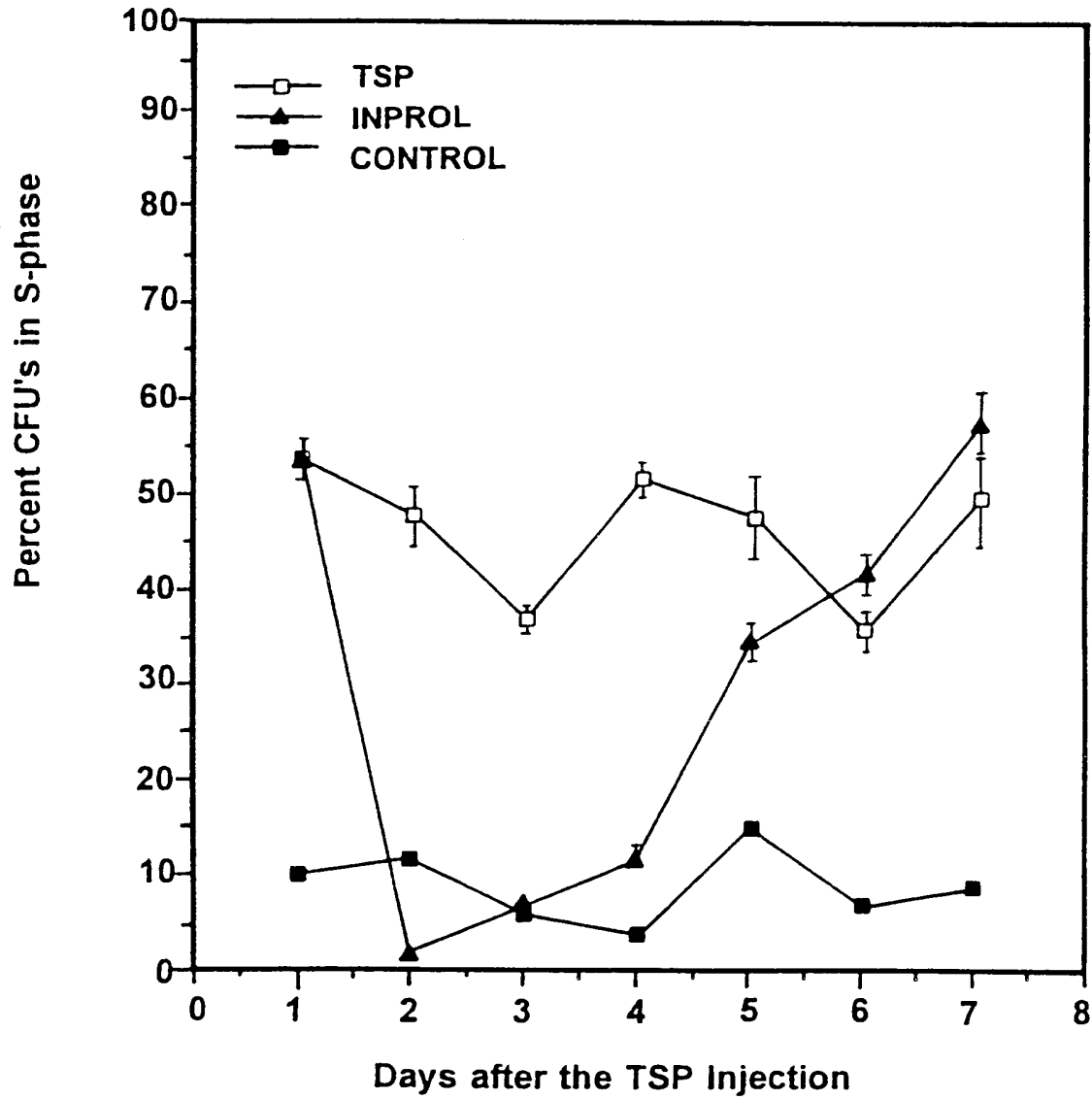
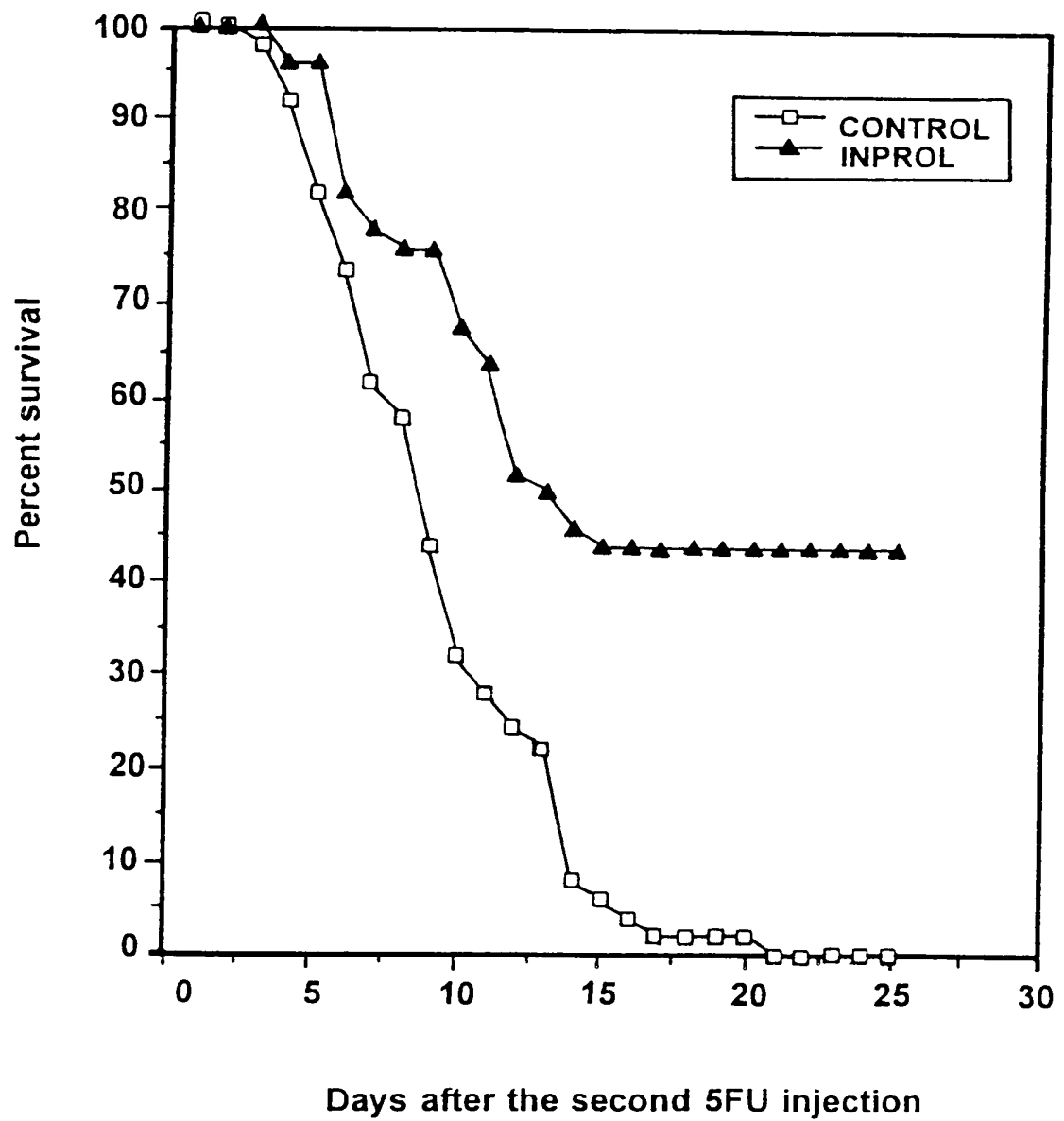


FIG. 7

FIG. 8



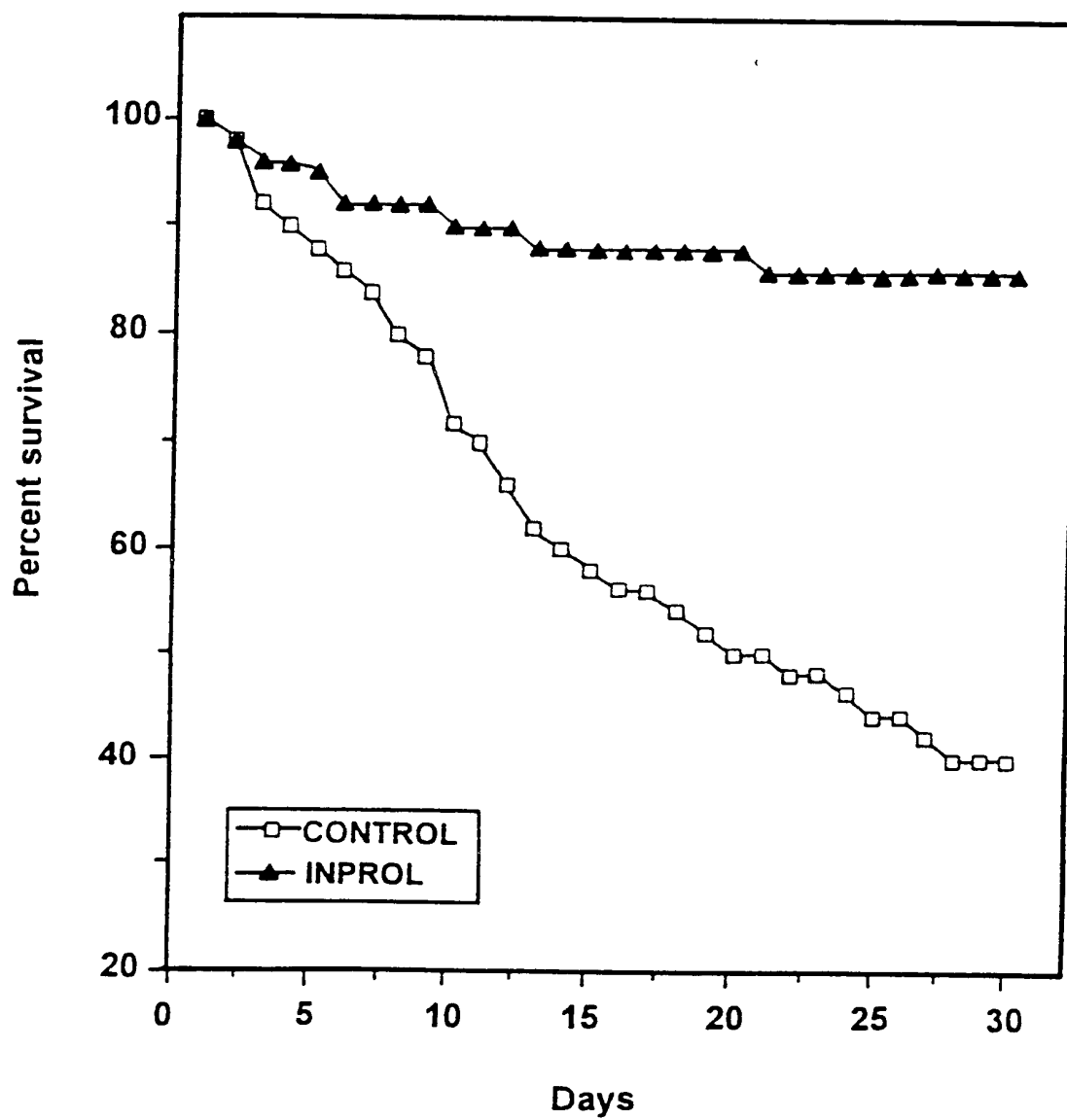


FIG. 9

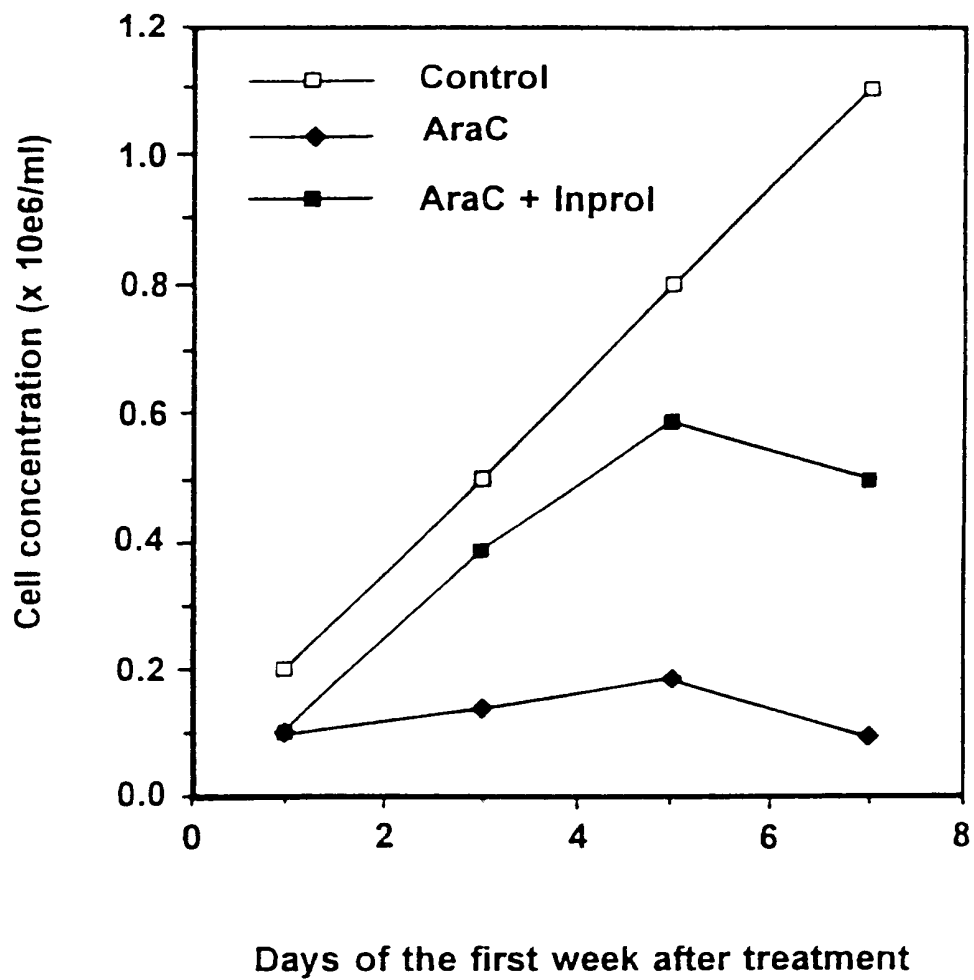


FIG. 10A

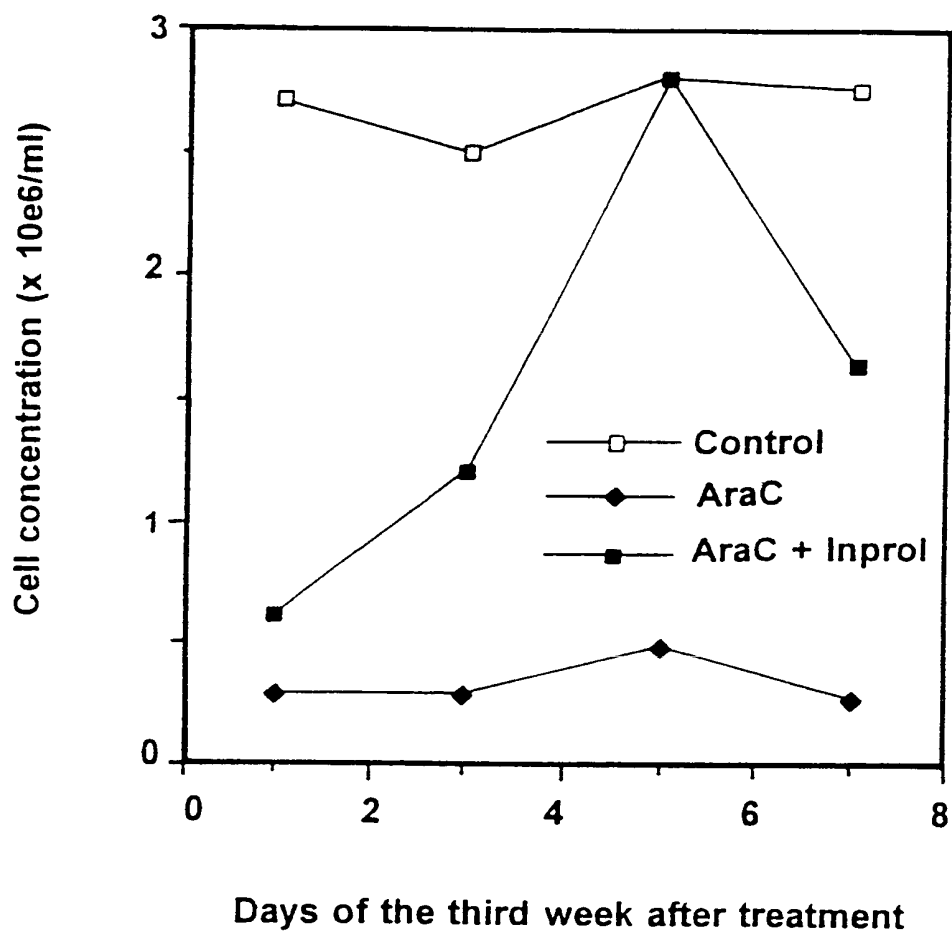


FIG. 10B

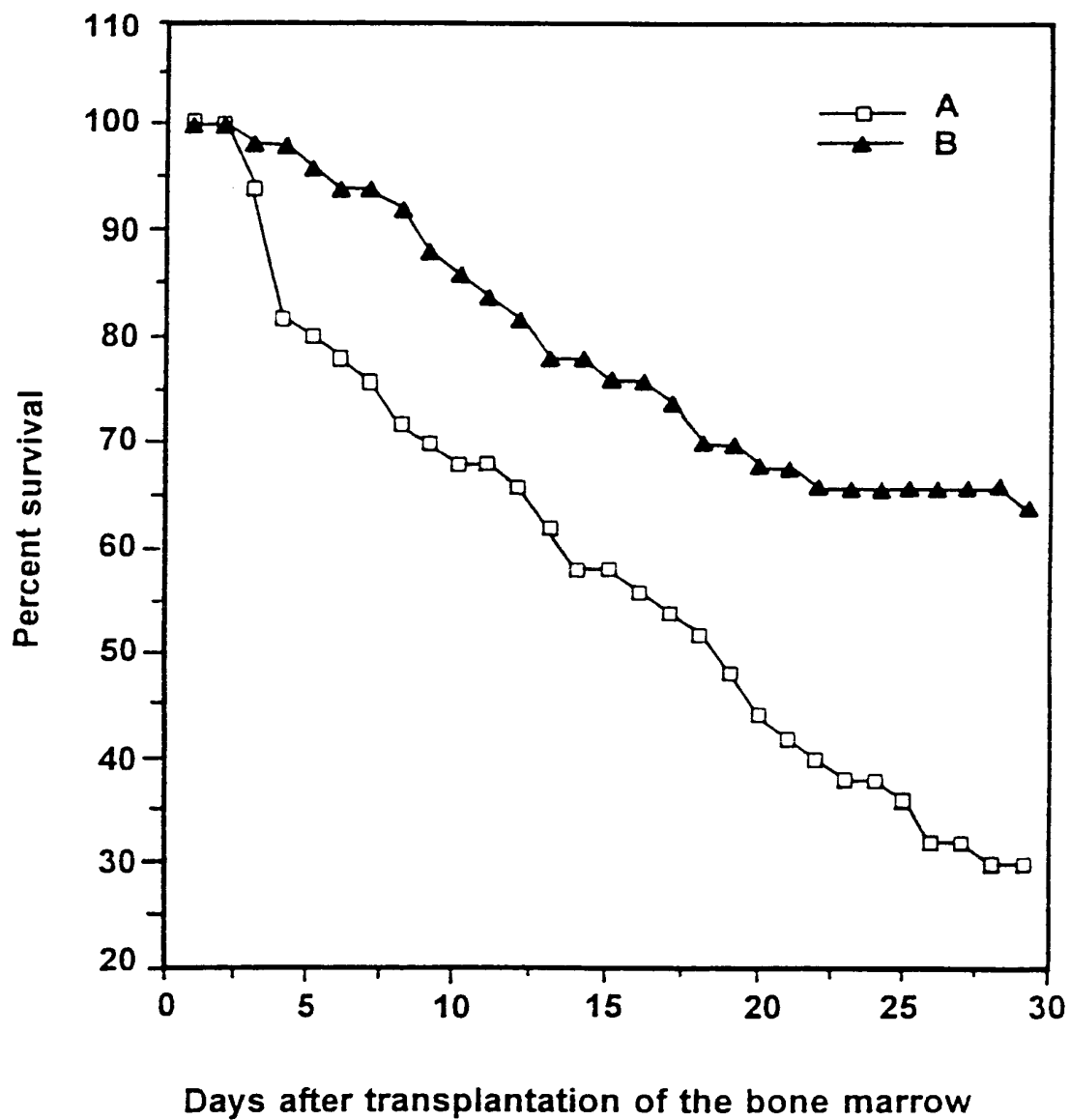


FIG. 11

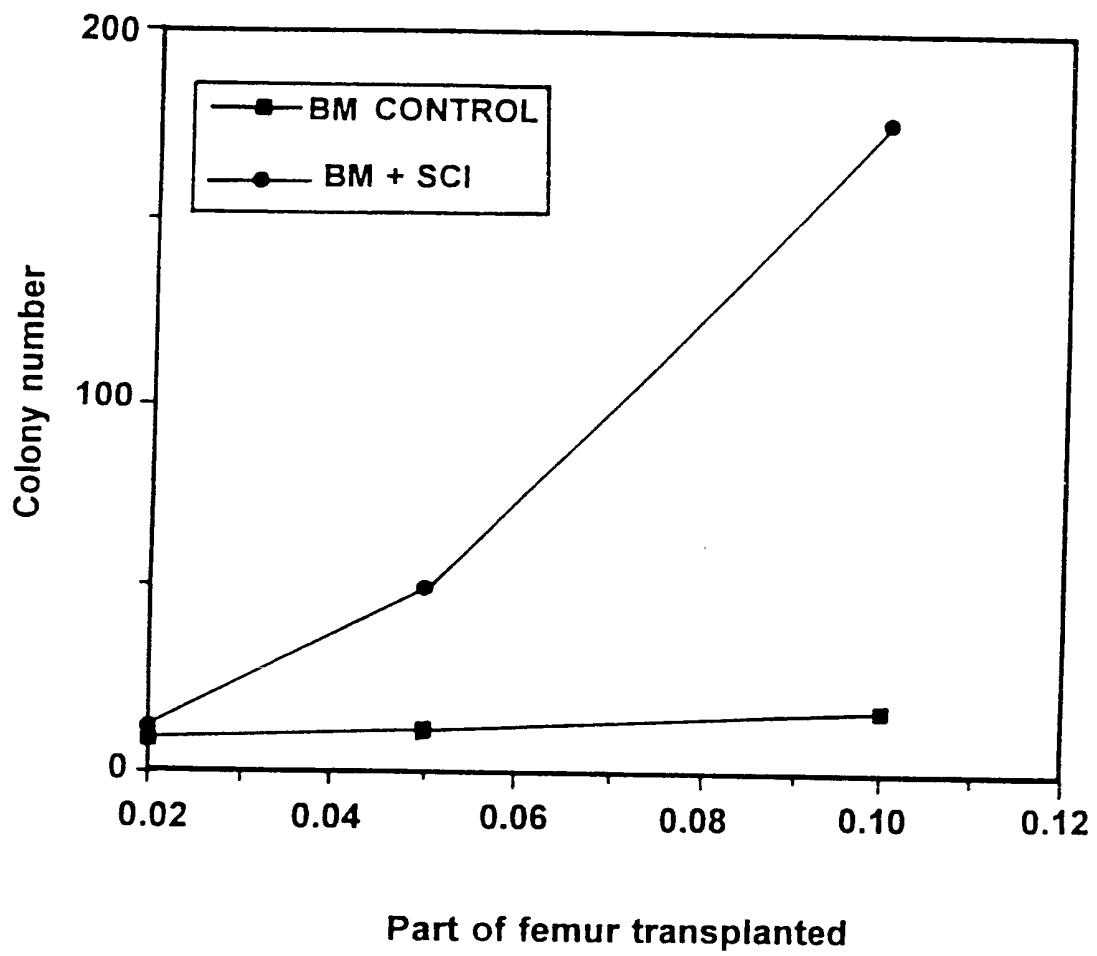


FIG. 12

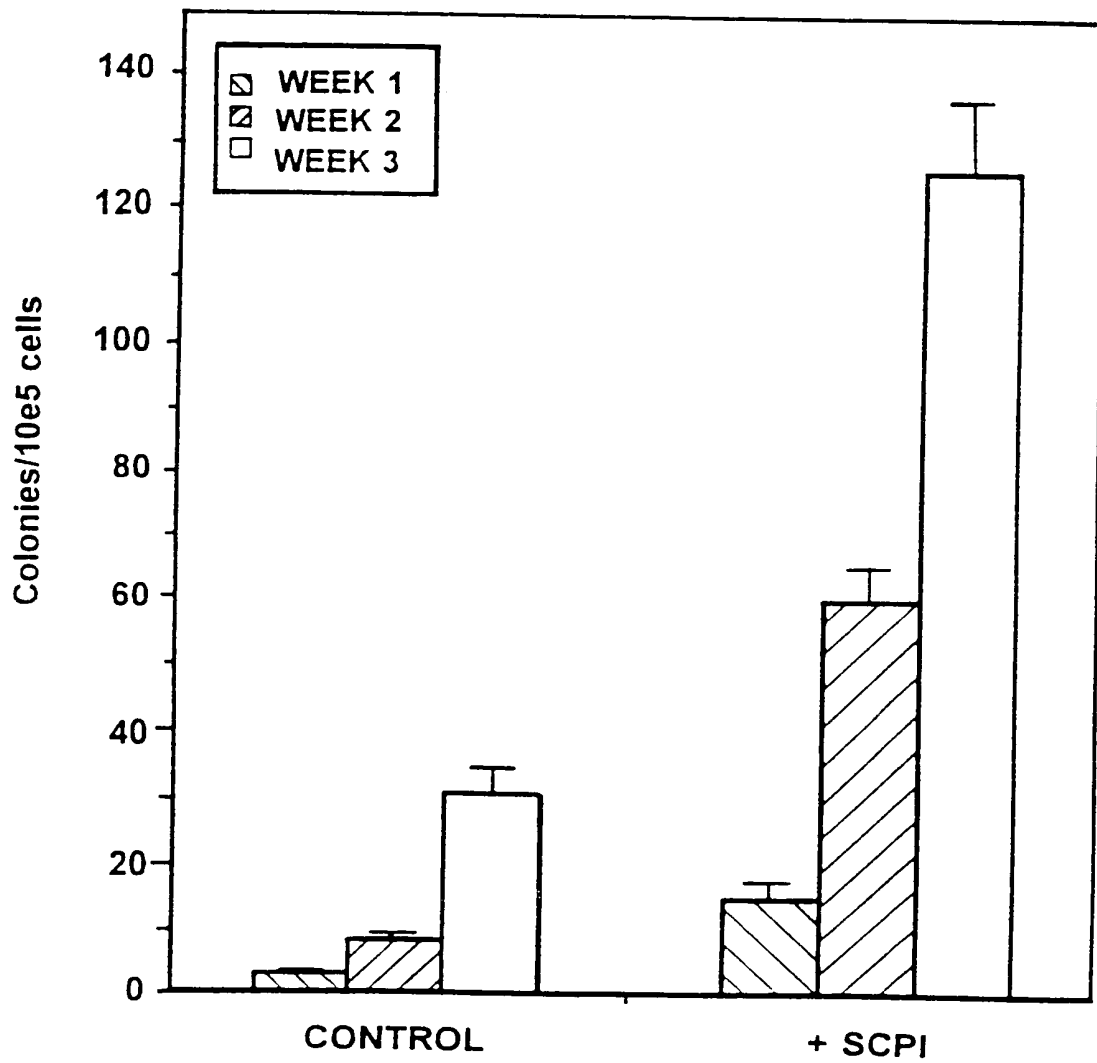


FIG. 13

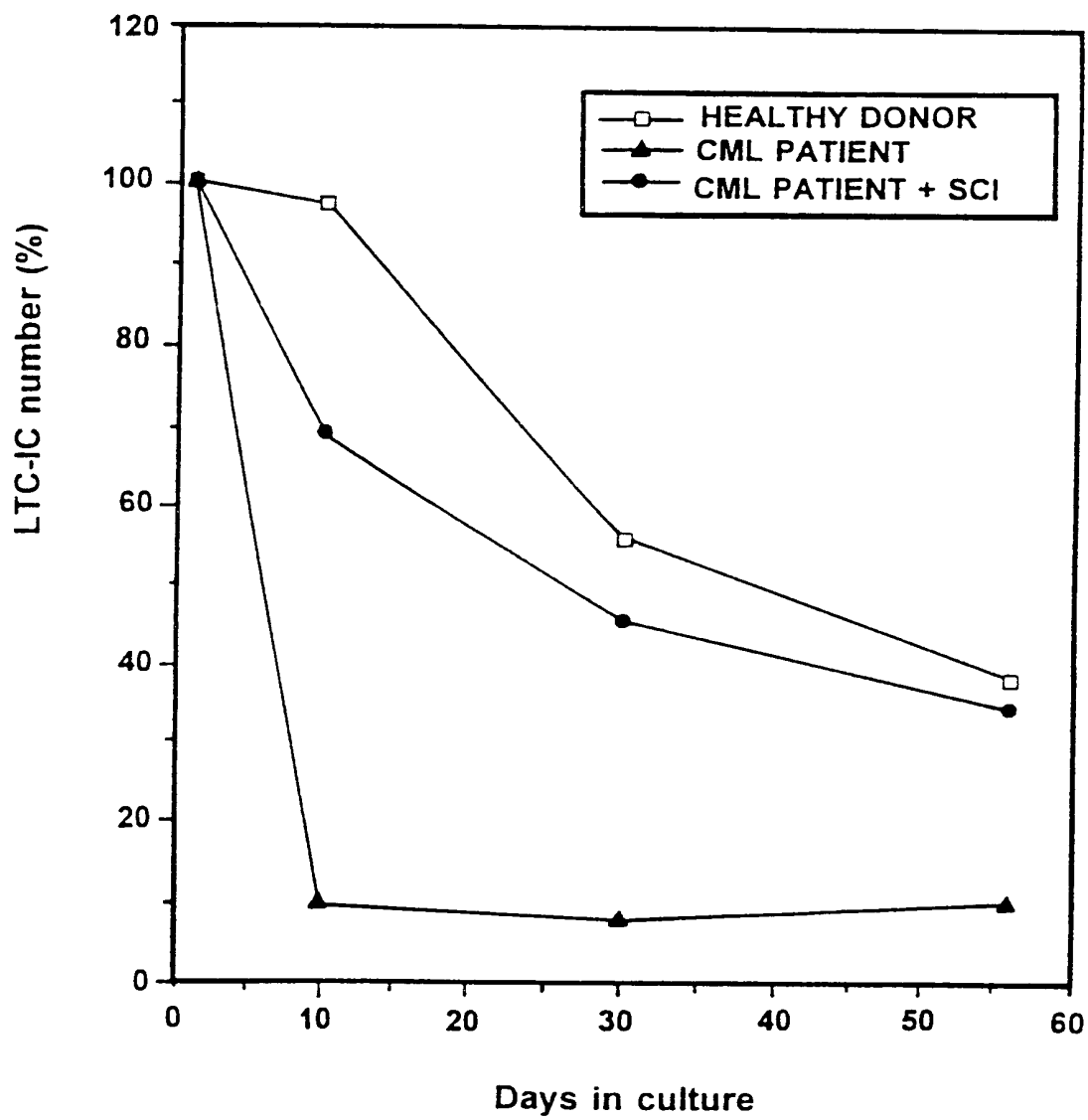
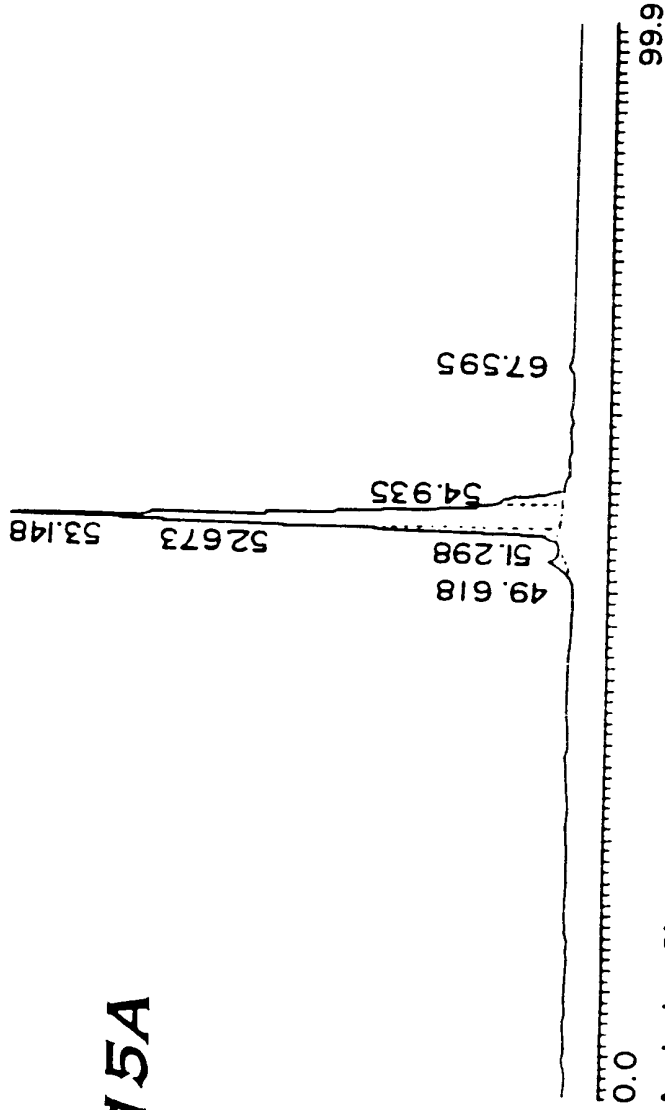


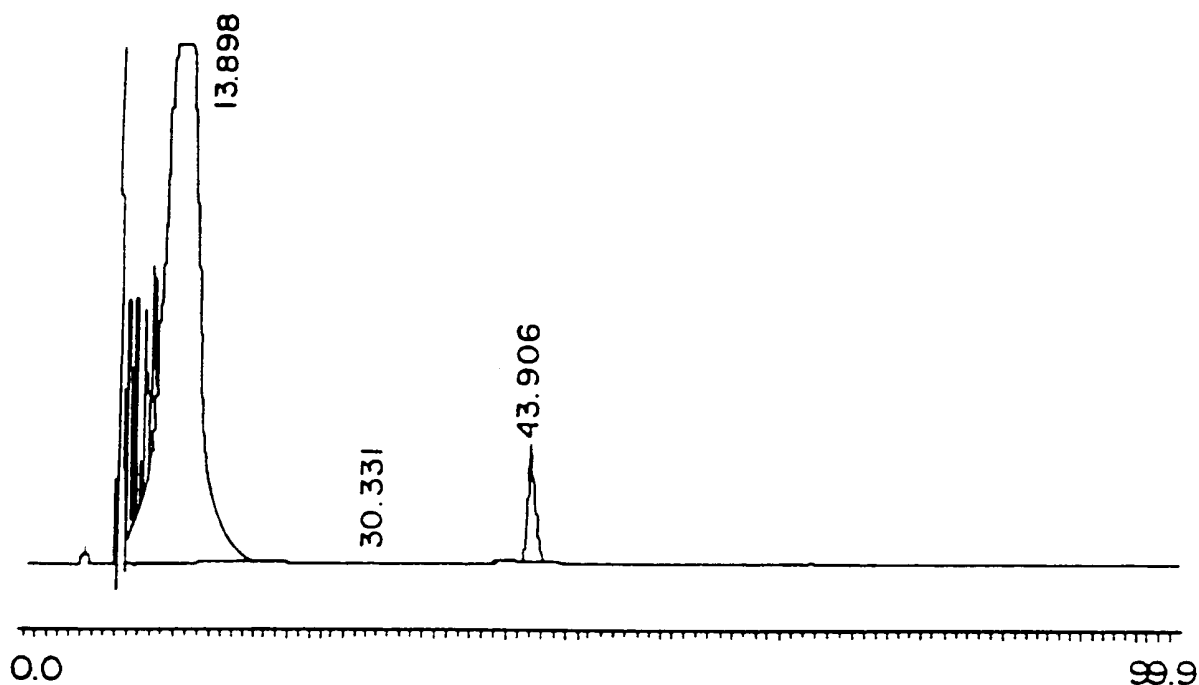
FIG. 14

FIG. 15A



Analysis: Channel A

Peak No.	Time	Type	Height(μY)	Area(μY-sec)	Area%
1	3.126	N1	691	7578	0.041
2	3.315	N2	1011	5150	0.027
3	49.618	N	8584	349227	1.893
4	51.298	N	1456	20274	0.109
5	52.673	N1	138069	2633395	14.278
6	53.148	N2	271587	14050458	76.181
	54.935	N3	33016	1332820	7.226
	67.595	N	3270	44507	0.241
TOTAL AREA				18443409	99.996



Analysis: Channel A

Peak No.	Time	Type	Height(μ Y)	Area(μ Y-sec)	Area%
1	4.383	N1	3945	95125	0.119
2	5.080	N2	28639	330889	0.413
3	5.216	N3	49084	531867	0.665
4	7.980	N1	399424	1110511	1.389
5	8.100	Err	1203320	2882013	3.605
6	8.241	N3	443249	1506159	1.884
7	8.386	N4	481563	2185702	2.734
8	8.533	N5	412886	1826165	2.284
9	8.701	N6	321500	842122	1.053
10	8.745	N7	404661	1610380	2.014
11	8.995	N8	435765	2489721	3.114
12	9.316	N9	517790	4801831	6.007

FIG. 15B

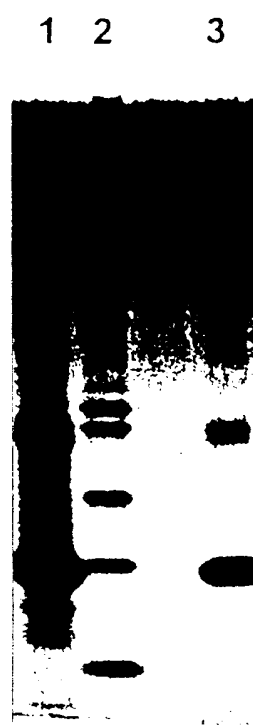


FIG. 15C

FIG. 16A

[illegible]

FIG. 16B

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20
 Val His Leu Thr Pro Glu Glu Lys Ser Ala Val Thr Ala Leu Trp Gly Lys Val Asn Val
 CIG CAC CTC ACT CCT CAG CAG AAC TCT GCC GTT ACT CCC CIG TGG GGT AAC GIG AAC GIG

 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40
 Asp Glu Val Gly Gly Glu Ala Leu Gly Arg Leu Leu Val Val Tyr Pro Trp Thr Gln Arg
 CAT GAA GTT GGT GAG GCC CIG GGC AGG CIG CIG GTC GTC TAC CTT TGG ACC CAG AGG

 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60
 Phe Phe Glu Ser Phe Gly Asp Leu Ser Thr Pro Asp Ala Val Met Gly Asn Pro Lys Val
 TTC TTT GAG TCC TTT GCG CAT CIG TCC ACT CCT CAT GCT GTT ATG GGC AAC CTT AAG CIG

 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80
 Lys Ala His Gly Lys Lys Val Leu Gly Ala Phe Ser Asp Gly Leu Ala His Leu Asp Asn
 AAG CTT CAT GCC AAG AAA GTC CTC GGT CCC TTT ACT GAT GGC CIG GCT CAC CIG CAC AAC

 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100
 Leu Lys Gly Thr Phe Ala Thr Leu Ser Glu Leu His Cys Asp Lys Leu His Val Asp Pro
 CIG AAG GGC ACC TTT GCC ACA CIG ACT GAG CIG CAC TGT CAC AAC CIG CAC GTC CAT CTT

 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120
 Glu Asn Phe Arg Leu Leu Gly Asn Val Leu Val Cys Val Leu Ala His His Phe Gly Lys
 GAG AAC TTC ACC CIG CIG GGC AAC GTC CIG GTC TGT GTC CIG GTC CAT CAC TTT GGC AAA

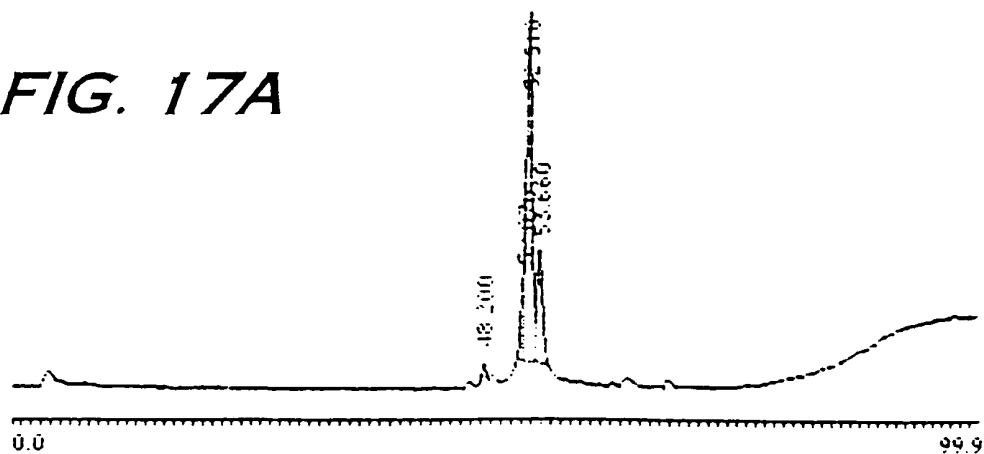
 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140
 Glu Phe Thr Pro Val Gln Ala Ala Tyr Gln Lys Val Val Ala Gly Val Ala Asn Ala
 GAA TTC ACC CCA CCA CIG CAG GCT GCC TAT CAG AAA GTC GTC GCT GGT GTC GCT AAT GCT

 141 142 143 144 145 146
 Leu Ala His Lys Tyr His
 CIG GCC CAC AAG TAT CAC

FIG. 16C

hHemA . pep	1	V LSPADKIN	20	VRAAKGKVG	30	HA GEYGAE	40	LE RFLSFF	50	TT KTVPFH
hHemB . pep	1	VHLTPPEKSA		VFL LIGKV		-NVDEVGSEA		LG RLLAVYF		WTORFESFG
mHemA . pep	1	VLSIEDKSN		IKRAATKIGG		HG-AEYGAE		LE RFEASEE		TT KTVPFH
mHemB . pep	1	VHLIDAEKAA		VSC LIGKVLS		E---EVGSEA		L GRLLVYF		WTORFESFG
pHemA . pep	1	VLSPADKAN		VRAAKGKVG		QA GAHGAER		LE RFLGFEF		TT KTVPFH
pHemB . pep	1	VHLSAEKSA		VGL LIGKVW		E---EVGSEA		L GRLLVYF		WTORFESFG
hHemA . pep	51	DLSH-----G	60	SAQVKSHGKK	70	VADALIT	80	A AHVDDMEN	90	ALS--ALS
hHemB . pep	51	DLSTPDVAVG		NPKVKAHGKK		VGA---ESP		GLAHLDNKKG		TFA--TLSL
mHemA . pep	51	DVSH-----G		SAQVKSHGKK		VADALAS		AGHLDDLEG		ALS--ALS
mHemB . pep	51	DLSSASALNG		NA SVKAHGKK		V---ITFED		GLNHLDSKKG		TEASL--SEL
pHemA . pep	51	NLSH-----G		SDQVKAHGKK		VADALIK		AVGHLDLPG		ALS--ALS
pHemB . pep	51	DLSTPDVAVG		NPKVKAHGKK		V---LQSFSD		GLKHLDNKKG		TFAKL--SEL
hHemA . pep	101	HAKKLAVDPE	110	NFKLLSHCLL	120	VTLAAHLPAS	130	ETPAVVASL	140	-KFLASVSTV
hHemB . pep	101	HCDKLAVDPE		NFRLLGNVLY		CVLAHFGGE		ETPFPVQAAQ		-KVAAGVAIA
mHemA . pep	101	HAKKLAVDPE		NFKLLSHCLL		VTLASHIPAD		ETPAVVASL		-KFLASVSTV
mHemB . pep	101	HCDKLAVDPE		NFRLLGNVIV		IVLGHILGKD		ETPAQAAT		QKVAAGVATA
pHemA . pep	101	HAKKLAVDPE		NFKLLSHCLL		VTLAAHHPDC		ETPSPVSLD		-KFLASVSTV
pHemB . pep	101	HCDKLAVDPE		NFRLLGNVIV		VTLARLGLHD		ETPDVQAA		QKVAAGVAIA
hHemA . pep	151	LISKVR	160		170	180	190	200		
hHemB . pep	151	LAHKYI								
mHemA . pep	151	LISKVR								
mHemB . pep	151	LAHKYI								
pHemA . pep	151	LISKVR								
pHemB . pep	151	LAHKYI								

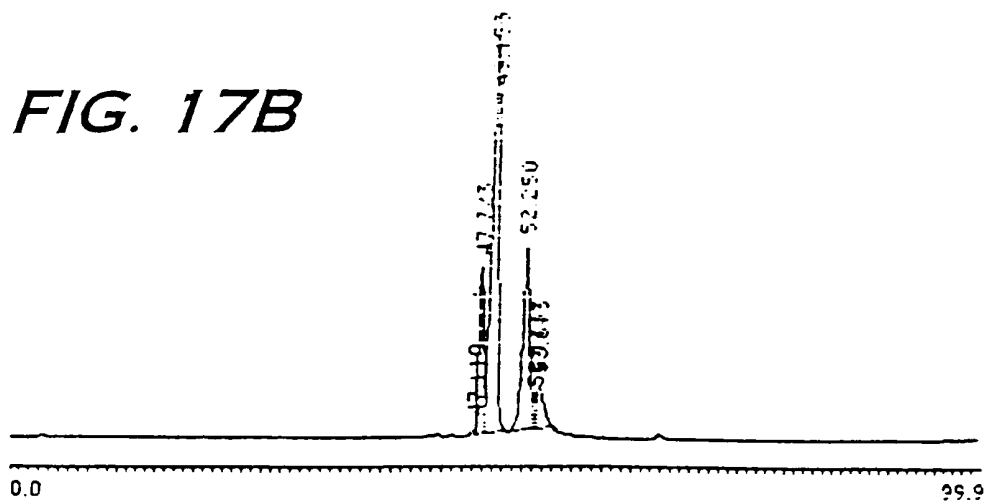
FIG. 17A



Analysis: Channel A

Peak No.	Time	Type	Height(μV)	Area(μV-sec)	Area%
1	48.200	N	1677	20438	1.515
2	52.076	N1	2625	116393	8.631
3	52.510	N2	32010	881490	65.362
4	53.660	N3	10066	330153	24.483
Total Area				1348474	99.996

FIG. 17B



Analysis: Channel A

Peak No.	Time	Type	Height(μV)	Area(μV-sec)	Area%
1	47.110	N1	1727	24840	0.204
2	47.723	N2	75067	1738939	14.321
3	49.153	N3	188795	6206410	51.114
4	52.250	N1	81476	3046748	25.092
5	53.113	N2	13195	202166	1.664
6	53.613	N3	19211	914954	7.535
	65.753	N	813	8066	0.066
Total Area				12142123	99.996

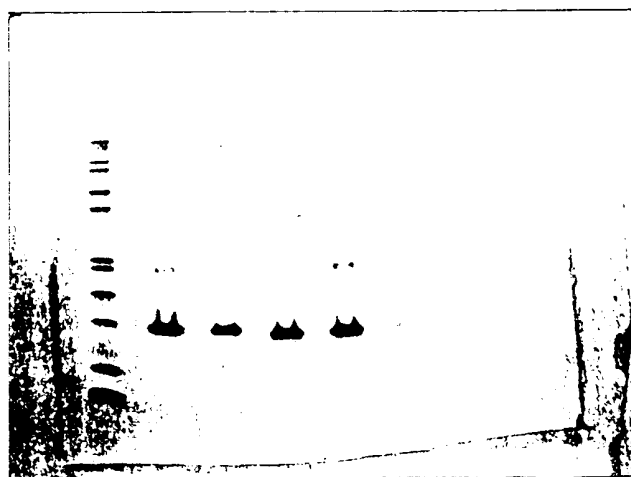


FIG. 18

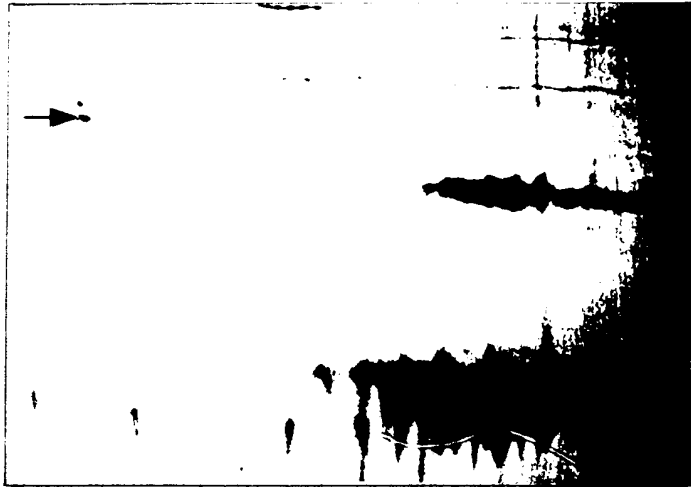


FIG. 19A

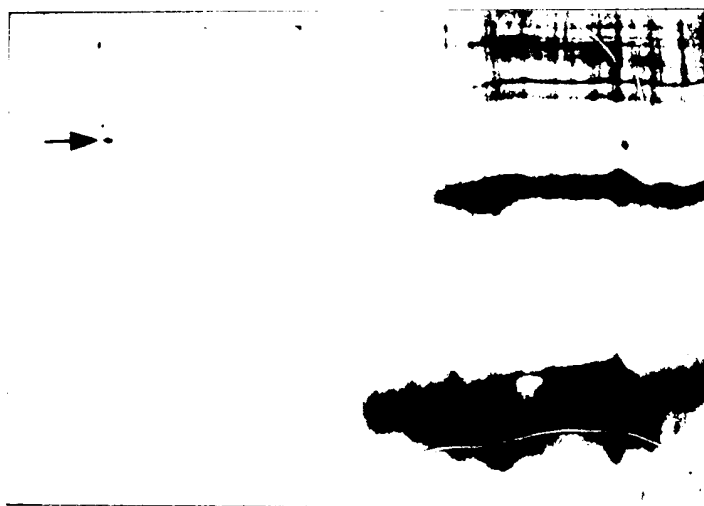


FIG. 19B

FIG. 20

